



PCT/JP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/049,957

DATE: 08/30/2002
TIME: 15:40:25

Input Set : A:\KATO21.txt
Output Set: N:\CRF4\08302002\J049957.raw

3 <110> APPLICANT: KATO, Yukio
4 FUJIMOTO, Katsumi
6 <120> TITLE OF INVENTION: CHONDROGENESIS PROMOTERS
8 <130> FILE REFERENCE: KATO-21
10 <140> CURRENT APPLICATION NUMBER: 10/049,957
11 <141> CURRENT FILING DATE: 2002-02-19
13 <150> PRIOR APPLICATION NUMBER: PCT/JS00/05590
14 <151> PRIOR FILING DATE: 2000-08-21
16 <150> PRIOR APPLICATION NUMBER: JP 232966/1999
17 <151> PRIOR FILING DATE: 1999-08-19
19 <160> NUMBER OF SEQ ID NOS: 15
21 <170> SOFTWARE: PatentIn version 3.1
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 2388
25 <212> TYPE: DNA
26 <213> ORGANISM: Oryctolagus cuniculus
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33 tggagggtgc gtgtgtcaccgc ggcgtccggcgc cccaggcaggca 180
35 aggccgttcgc cggacggccgc ctccacggcc cccctgtgtgc cttgtcggccg 240
37 accactcggtt ccacgtccatc ggcggcccaaggc acggccaggc 300
39 ccatttacgca ggccggaaag gaaacacggcc tgaaggccgt gttggccgaa 360
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47 tcaaaaggccgtt cggccgtatc ttccggggaa gctgtgttgc cccgttcc 600
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51 agaccccccgtt ggagccgttac taacgttaca cccggggccctt cccgtgtccgttgc 720
53 caggggacatgtt ggccttgcgtt aacccacacgc cccggccatc 780
55 tgcctccgtt gggccacatgtt cttgtgttgc cccggccatc 840
57 gccggccatc cgttccacccgtt gggacttgc gttgtgttgc cccggccatc 900
59 ttgggtgtccgc ggccggacaccgc gacggccaggc tccatcttcgc gtttctcaat 960
61 ggctgttgc ccacggaggc agoagcttgc agatgttgc cccggggcc 1020
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69 acatccgttgc ggccttgcgtt cggccggaggc gacccggccatc 1260
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75 tggatgttgcggcc gacccggaggc agtacactgttgc acgttgcgttgc 1440
77 ggccttgcgttgc cccggccatc acatccgttgc tccatcttcgc 1500
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85	ggccaaacaa	gtgactggc	aaacgcccgg	agcggtaacta	tgccgacagt	ggccgccttc	1740										
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89	acacaaatgg	ccacaatccc	gacccgtggg	ctgcccatact	gaggagccag	gactacgagc	1860										
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93	agattccgtc	ccacggccgtc	atgggtcgcc	ccgacacccaa	catttttcacc	gtttacggac	1980										
95	tgctggacaa	ggcccgaggc	ctgtttgggg	agcaccacaa	caagaacggg	ttcaagatgt	2040										
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124						35				40				45			
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133						85				90				95			
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136						100				105				110			
138	Val	Lys	Arg	Ser	Ser	Asn	Val	Thr	Ile	Asn	Thr	Leu	Arg	Gly	Val	Lys	
139						115				120				125			
141	Ser	Cys	His	Thr	Gly	Ile	Asn	Arg	Thr	Val	Gly	Trp	Asn	Val	Pro	Val	
142						130				135				140			
144	Gly	Tyr	Leu	Val	Asp	Ser	Gly	Arg	Leu	Ser	Val	Met	Gly	Cys	Asp	Val	
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147	Leu	Lys	Ala	Val	Ser	Glu	Tyr	Phe	Gly	Gly	Ser	Cys	Val	Pro	Gly	Ala	
148						165				170				175			
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151						180				185				190			
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157						210				215				220			
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162 Leu Pro Ser Trp Gly His Met Leu Met Ser Arg Asp Phe Glu Leu Leu
163 245 250 255
165 Cys Arg Asp Gly Ser Arg Ala Ser Val Thr Glu Trp Gln His Cys His
166 260 265 270
168 Leu Ala Arg Val Pro Ala His Ala Val Val Val Arg Ala Asp Thr Asp
169 275 280 285
171 Ala Gly Leu Ile Phe Arg Leu Leu Asn Glu Gly Gln Arg Leu Phe Ser
172 290 295 300
174 His Glu Gly Ser Ser Phe Gln Met Phe Ser Ser Glu Ala Tyr Gly Gln
175 305 310 315 320
177 Lys Asn Leu Leu Phe Lys Asp Ser Thr Leu Glu Leu Val Pro Ile Ala
178 325 330 335
180 Thr Gln Thr Tyr Glu Ala Trp Leu Gly Pro Glu Tyr Leu His Ala Met
181 340 345 350
183 Lys Gly Leu Leu Cys Asp Pro Asn Arg Leu Pro Pro Tyr Leu Arg Trp
184 355 360 365
186 Cys Val Leu Ser Thr Pro Glu Ile Gln Lys Cys Gly Asp Met Ala Val
187 370 375 380
189 Ala Phe Ser Arg Gln Arg Leu Lys Pro Glu Ile Gln Cys Val Ser Ala
190 385 390 395 400
192 Glu Ser Pro Gln His Cys Met Glu Gln Ile Gln Ala Gly His Ile Asp
193 405 410 415
195 Ala Val Thr Leu Asn Gly Glu Asp Ile His Thr Ala Gly Lys Thr Tyr
196 420 425 430
198 Gly Leu Ile Pro Ala Ala Gly Glu Leu Tyr Ala Ala Asp Asp Arg Ser
199 435 440 445
201 Asn Ser Tyr Phe Val Val Ala Val Val Lys Arg Asp Ser Ala Tyr Ala
202 450 455 460
204 Phe Thr Val Asp Glu Leu Arg Gly Lys Arg Ser Cys His Pro Gly Phe
205 465 470 475 480
207 Gly Ser Pro Ala Gly Trp Asp Val Pro Val Gly Ala Leu Ile His Trp
208 485 490 495
210 Gly Tyr Ile Arg Pro Arg Asn Cys Asp Val Leu Thr Ala Val Gly Gln
211 500 505 510
213 Phe Phe Asn Ala Ser Cys Val Pro Val Asn Asn Pro Lys Lys Tyr Pro
214 515 520 525
216 Ser Ser Leu Cys Ala Leu Cys Val Gly Asp Glu Gln Gly Arg Asn Lys
217 530 535 540
219 Cys Thr Gly Asn Ser Gln Glu Arg Tyr Tyr Gly Asp Ser Gly Ala Phe
220 545 550 555 560
222 Arg Cys Leu Val Glu Gly Ala Gly Asp Val Ala Phe Val Lys His Thr
223 565 570 575
225 Thr Ile Phe Asp Asn Thr Asn Gly His Asn Pro Glu Pro Trp Ala Ala
226 580 585 590
228 His Leu Arg Ser Gln Asp Tyr Glu Leu Leu Cys Pro Asn Gly Ala Arg
229 595 600 605
231 Ala Glu Ala His Gln Phe Ala Ala Cys Asn Leu Ala Gln Ile Pro Ser
232 610 615 620
234 His Ala Val Met Val Arg Pro Asp Thr Asn Ile Phe Thr Val Tyr Gly

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235	625	630	635	640													
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240	Gly	Phe	Lys	Met	Phe	Asp	Ser	Ser	Ser	Tyr	His	Gly	Arg	Asp	Leu	Leu	
241																	670
243	Phe	Lys	Asp	Ala	Thr	Val	Arg	Ala	Val	Pro	Val	Gly	Glu	Arg	Thr	Thr	
244																	685
246	Tyr	Gln	Asp	Trp	Leu	Gly	Pro	Asp	Tyr	Val	Ala	Ala	Leu	Glu	Gly	Met	
247																	700
249	Gln	Ser	Gln	Arg	Cys	Ser	Gly	Ala	Ala	Val	Gly	Ala	Pro	Gly	Ala	Ser	
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266	atggagggtgc	tggtgtgcgc	cacccggac	ccagacggc	caaaatgcgg	caacatgcgg	caacatgcgg	180									
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270	gaccactg	tcggatctat	cgcggccccc	gaggctgcgg	ccatccatct	ggatggaggg	ggatggaggg	300									
272	gccccatctat	aggcgggaaa	ggagcacggc	ctgaaggccg	tggggccgca	ttggggccgca	ttggggccgca										
274	caaaatgcgt	gtatccatctt	ttacggccgt	tgctgtgtgc	ggggaggccgc	ccatgtgcac	ccatgtgcac	420									
276	atggacaccc	ttggaaatgg	gtggatctgc	cacacggca	tcaatgcac	atgtggctgg	atgtggctgg										
278	aacgtgcgg	tggggatct	gtgtggatgc	ggccggctct	cggtgtatgg	ctggatgtta	ctggatgtta	540									
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318	ggccgcacatgc	atgtgtatgc	aaatgtatgc	ggatgtatgc	ttactatgtatgc	tgccgtgtatgc	tgccgtgtatgc										
320	atgtgtatgc	tggggatataatgg	gggtgtatgc	gggtgtatgc	gttgcgtatgc	ggccacacatgc	ggccacacatgc										
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338	cggccgcggcc	gccccagage	tccgatggcc	gcccggggag	tttccgcgc	ggccctctcgcc	2340
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356	Gln His Lys Cys Gly Asn Met Ser Glu Ala Phe Arg Glu Ala Gly Ile						
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359	Gln Pro Ser Leu Leu Cys Val Arg Gly Thr Ser Ala Asp His Cys Val						
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362	Gln Leu Ile Ala Ala Gln Glu Ala Asp Ala Ile Thr Leu Asp Gly Gly						
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377	Gly Tyr Leu Val Glu Ser Gly Arg Leu Ser Val Met Gly Cys Asp Val						
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380	Leu Lys Ala Val Ser Asp Tyr Phe Gly Gly Ser Cys Val Pro Gly Ala						
381		165	170			175	
383	Gly Glu Thr Ser Tyr Ser Glu Ser Leu Cys Arg Leu Cys Arg Gly Asp						
384		180	185			190	
386	Ser Ser Gly Glu Gly Val Cys Asp Lys Ser Pro Leu Glu Arg Tyr Tyr						
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392	Ala Phe Val Lys His Ser Thr Val Leu Glu Asn Thr Asp Gly Lys Thr						
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VERIFICATION SUMMARY

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